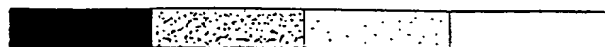


FIG.1

Fas/APO1	VKGFRKNGVNEAKIDEIKNDNVQDTAEQKVQLLRNWHQHLHGKKEA.YDILIKDLKKANLCTHA	307
Ankyrin1	WAEAREIQFSVEDINRIRVENPNSLIEQSVALLNIMVIREGQN.ANMENLYTALQSIDRGEIV	1479
p55 TNF-R	WKEFVRRLGLSDHEIDRLQLNGRQLREAQYSMLATWRRRTPRREATLELLGRVLRDHDLLGCL	434
NGF-R	WRHLAGELGYPPEHIDSFTHIE....ACPVRLLASWA...TQDSATLDALLAALRRRIQRADLV	414
MORT1	WRLARQLKVSDTKIDSIEDRYPRNLIERVRESLRIMKN.TEKENATVAHLVGAALRSCQM.LV	210

FIG. 2

DBD CONSTRUCT	AD CONSTRUCT	SCORE	lpr <sup>cg</sup> -like mutation effect		
			IN DBD CONSTRUCT	IN AD CONSTRUCT	IN BOTH
p55	p55				
Fas	Fas				
TRADD	TRADD		ND	ND	ND
MORT1	MORT1				
RIP	RIP				
TRADD	p55		ND		ND
p55	TRADD			ND	ND
TRADD	Fas		ND		ND
Fas	TRADD			ND	ND
MORT1	p55				
p55	MORT1				
MORT1	Fas				
Fas	MORT1				
RIP	p55				
p55	RIP				
RIP	Fas				
Fas	RIP				
TRADD	MORT1		ND		ND
MORT1	TRADD			ND	ND
TRADD	RIP		ND		ND
RIP	TRADD			ND	ND
MORT1	RIP				
RIP	MORT1				



<1h      1-2h      2-24h      NEGATIVE

204070 204500

FIG. 3

